

In the claims:

Please amend claims 8, 35, 36, 38, 39, 41, 42, and 44 as follows: (For the Examiner's convenience, all of the pending claims, whether or not amended, are reproduced below.)

D1  
1. ~~8.~~ (Twice Amended) An isolated nucleic acid comprising the nucleotide sequence shown in [the Sequence Listing] SEQ ID NO:1 or a fragment of the nucleotide sequence shown in [the Sequence Listing] SEQ ID NO:1 which encodes an antigenic fragment of PM-1 protein.

2. ~~9.~~ The nucleic acid of claim 8, which is cDNA.

D2  
3. ~~35.~~ (Amended) The nucleic acid of claim 8 wherein the nucleotide sequence comprises the coding region of the nucleotide sequence shown in [the Sequence Listing] SEQ ID NO:1.

4. ~~36.~~ (Amended) The nucleic acid of claim 35, wherein the coding region comprises nucleotide 179 to nucleotide 1627 of the nucleotide sequence shown in [the Sequence Listing] SEQ ID NO:1.

5. ~~37.~~ A nucleic acid comprising a nucleotide sequence which is a functional equivalent of the nucleic acid of claim 8.

D3  
6. ~~38.~~ (Amended) The nucleic acid of claim 37, wherein the nucleotide sequence hybridizes to a nucleotide sequence which is complementary to the nucleotide sequence shown in [the Sequence Listing] SEQ ID NO:1.

D3

<sup>7</sup> 39. (Amended) An isolated nucleic acid comprising a nucleotide sequence which encodes the amino acid sequence shown in [the Sequence Listing] SEQ ID NO:1 or a nucleotide sequence which encodes an antigenic fragment of the amino acid sequence shown in [the Sequence Listing] SEQ ID NO:1.

<sup>8</sup> 40. A nucleic acid comprising a nucleotide sequence which is a functional equivalent of the nucleic acid of claim <sup>7</sup> 39.

D4

<sup>9</sup> 41. (Amended) The nucleic acid of claim <sup>8</sup> 40, wherein the nucleotide sequence hybridizes to a nucleotide sequence which is complementary to the nucleotide sequence shown in [the Sequence Listing] SEQ ID NO:1.

<sup>10</sup> 42. (Amended) The nucleic acid of claim <sup>7</sup> 39, wherein the antigenic fragment comprises at least one T cell epitope which is recognized by a T cell receptor specific for the PM-1 protein having an amino acid sequence shown in [the Sequence Listing] SEQ ID NO:1.

<sup>11</sup> 43. The nucleic acid of claim <sup>10</sup> 42, wherein the antigenic fragment comprises at least 7 amino acid residues.

<sup>12</sup> 44. (Amended) The nucleic acid of claim <sup>6</sup> 38, which encodes an amino acid sequence shown in [the Sequence Listing] SEQ ID NO:1 which is modified by an amino acid substitution, deletion, or addition.

D5

13  
45. A recombinant expression vector comprising the nucleic acid of claim 8.

14  
46. A recombinant expression vector comprising the nucleic acid of claim 35.

15  
47. A recombinant expression vector comprising the nucleic acid of claim 39.

16  
48. A recombinant expression vector comprising the nucleic acid of claim 42.

17  
49. A host cell transformed with the recombinant expression vector of claim

13  
45.

50. A host cell transformed with the recombinant expression vector of claim  
of claim 46.

51. A host cell transformed with the recombinant expression vector of claim  
of claim 47.

52. A host cell transformed with the recombinant expression vector of claim  
of claim 48.

#### REMARKS

In a telephone conversation with Examiner Scheiner on June 12, 1996, Applicants provisionally elected, with traverse, to prosecute the invention of Group II, claims 8, 9, and 35-52, Applicants also elected the species of the full-length nucleic acid sequence with encodes PM-1. Applicants hereby affirm this election.

U.S. Serial No.: 08/\_\_\_\_\_,605

-2-

Group Art Unit: 1813

<sup>18</sup>  
50. (Amended) A host cell transformed with the recombinant expression vector [of claim] of claim <sup>14</sup> 46.

<sup>19</sup>  
51. (Amended) A host cell transformed with the recombinant expression vector [of claim] of claim <sup>15</sup> 47.

<sup>20</sup>  
52. (Amended) A host cell transformed with the recombinant expression vector [of claim] of claim <sup>16</sup> 48.

#### REMARKS

Claims 8, 9 and 35-52 are pending in the application. Claims 50-52 have been amended. Any amendments to and/or cancellation of the claims should in no way be construed as an acquiescence to any of the objections and/or rejections of record. The amendments and/or cancellations are being made to expedite prosecution of the above-identified application. Applicants reserve the right to file the same or similar claims in this or another application. Claims 8, 9, 35, 37-41, 44-47 and 49-51 have been rejected under 35 USC §102(a).

#### *Rejection of Claims 8, 9, 35-41, 44-47 and 49-51 under 35 USC §102(a)*

Claims 8, 9, 35, 37-41, 44-47 and 49-51 were rejected under 35 USC §102(a) as being anticipated by Pietropaolo et al., *Diabetes* 40:1A, abstract #2.

The Examiner states:

Pietropaolo et al. teach the PM-1 protein wherein an initial sequence shows a 252bp open reading frame coding for 84 amino acids without significant homologies to known sequences. Pietropaolo et al. failed to disclose the specific nucleotide sequence of their clone. However, a sequence is merely a characterization of the DNA and Pietropaolo et al. teach that DNA which inherently possesses the claimed sequence. The vector and host cell are also taught.